(FILE 'HOME' ENTERED AT 14:14:49 ON 16 FEB 2000)

FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH' ENTERED AT 14:15:14 ON 16 FEB 2000

- L1 2226 S GALACTOSYL(W)TRANSFERASE
- L2 10756 S GALACTOSYLTRANSFERASE
- L3 12092 S L1 OR L2
- L4 1004 S FUCOSYL(W)TRANSFERASE
- L5 4276 S FUCOSYLTRANSFERASE
- L6 4747 S L4 OR L5
- L7 2001 S GLYCOSYL(W)TRANSFERASE
- L8 10329 S GLYCOSYLTRANSFERASE
- L9 11770 S L7 OR L8
- L10 3582 S HYPERACUTE(W)REJECTION
- L11 373146 S CHIMERIC OR HYBRID
- L12 602 S L3 AND L6
- L13 16 S L12 AND L11
- L14 0 S L13 AND L10
- L15 9 DUPLICATE REMOVE L13 (7 DUPLICATES REMOVED)
- L16 755308 S L11 OR FUSION
- L17 24 S L16 AND L12
- L18 13 DUPLICATE REMOVE L17 (11 DUPLICATES REMOVED)
- L19 7869 S CYTOPLASMIC(W)TAIL
- L20 108874 S SWITCHING OR SUBSTITUTING
- L21 3287155 S DOMAIN OR REGION
- L22 45815 S CATALYTIC(S)L21
- L23 67 S L19 AND L20 AND L21
- L24 25111 S L9 OR L5 OR L3
- L25 5 S L24 AND L23
- L26 1 DUPLICATE REMOVE L25 (4 DUPLICATES REMOVED)
- L27 163 S L19 AND L22
- L28 33 S L27 AND L24
- L29 14 S L28 AND L11
- L30 18 S L28 AND L16
- L31 4 DUPLICATE REMOVE L30 (14 DUPLICATES REMOVED)
- L32 33 S L27 AND L24
- L33 7 DUPLICATE REMOVE L32 (26 DUPLICATES REMOVED)
- L34 86 S L19 AND L24
- L35 39 S L34 AND L16
- L36 9 DUPLICATE REMOVE L35 (30 DUPLICATES REMOVED)
- L37 19 DUPLICATE REMOVE L34 (67 DUPLICATES REMOVED)
- L38 0 S SANDRIN?/AT
- L39 1731 S SANDRIN?/AU
- L40 10 S L39 AND L19
- L41 2 DUPLICATE REMOVE L40 (8 DUPLICATES REMOVED)
- L42 95 S L39 AND L24
- L43 42 S L42 AND L10
- L44 14 DUPLICATE REMOVE L43 (28 DUPLICATES REMOVED)
- L45 1755 S L16 AND L19
- L46 15523 S LOCALIZ?(P)GOLGI
- L47 124 S L46 AND L45
- L48 240516 S COMPETE OR OUTCOMPETE OR COMPETITION
- L49 14532 S L48(P)SUBSTRATE
- L50 1 S L49 AND L47
- L51 1 S L45 AND L49
- L52 3 S L48 AND L47

L53	3 DUPLICATE REMOVE L52 (0 DUPLICATES REMOVED)
L54	53 S CARBOHYDRATE(S)REMODELLING
L55	15 S L24 AND L54
L56	0 S L16 AND L55
L57	5 DUPLICATE REMOVE L55 (10 DUPLICATES REMOVED)

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Help	 }	
	Logout	
Main Menu Search Form Posting Counts	Show S Numbers	Edit S Numbers

3

150 and 126

Database: All Data	pases (USPT + EPAB + JPAB + DWPI + TDBD)	▼
Refine Search:	150 and 126	

Search History

DB Name	Query	Hit Count	Set Name
ALL	150 and 126	3	<u>L52</u>
ALL	150 and redirect	5	<u>L51</u>
ALL	149 and substrate	84	<u>L50</u>
ALL	147 and 148	100	<u>L49</u>
ALL	compete or outcompete or competition	35741	<u>L48</u>
ALL	124 and 132	209	<u>L47</u>
ALL	144 and 113	2	<u>L46</u>
ALL	114 and 144	1	<u>L45</u>
ALL	141 and 132	53	<u>L44</u>
ALL	114 and 142	1	<u>L43</u>
ALL	141 and chimeric	27	<u>L42</u>
ALL	hyperacute adj1 rejection	133	<u>L41</u>
ALL	139 and 114	16	<u>L40</u>
ALL	136 and 132	351	<u>L39</u>
ALL	137 and 132	5	<u>L38</u>
ALL	redirect and 136	10	<u>L37</u>
ALL	golgi and localiz\$	562	<u>L36</u>
ALL	134 and localiz\$	13	<u>L35</u>

ALL	133 and target\$	28	<u>L34</u>
ALL	114 and 132	43	<u>L33</u>
ALL	chimeric or hybrid	97552	<u>L32</u>
ALL	130 and chimeric	5	<u>L31</u>
ALL	126 and 128	40	<u>L30</u>
ALL	126 same 128	1	<u>L29</u>
ALL	reducing with carbohydrate	1056	<u>L28</u>
ALL	126 and 125	8	<u>L27</u>
ALL	18 or 113 or 110	896	<u>L26</u>
ALL	124 and 115	195	<u>L25</u>
ALL	119 and 123	280	<u>L24</u>
ALL	120 or 121 or 122	908	<u>L23</u>
ALL	cytoplasmic adj 1 tail	391	<u>L22</u>
ALL	amino-terminal adj 1 tail	1	<u>L21</u>
ALL	terminal adj1 tail	544	<u>L20</u>
ALL	117 or 118	2217	<u>L19</u>
ALL	cytoplasmic adj1 domain	1346	<u>L18</u>
ALL	catalytic adj1 domain	1020	<u>L17</u>
ALL	114 and 115	34	<u>L16</u>
ALL	chimeric or heterologous	18731	<u>L15</u>
ALL	113 and 110	114	<u>L14</u>
ALL	13 or 112	680	<u>L13</u>
ALL	glycosyl\$transferase	510	<u>L12</u>
ALL	galactosyl\$transferase	223	<u>L11</u>
ALL	14 or 19	218	<u>L10</u>
ALL	fucosyl\$transferase	205	<u>L9</u>
ALL	15 or 16	235	<u>L8</u>
ALL	15 and 16	220	<u>L7</u>
ALL	galactosyl\$transferase	223	<u>L6</u>
ALL	galactosyltransferase	232	<u>L5</u>
ALL	fucosyltransferase	216	<u>L4</u>
ALL	glycosyltransferase	662	<u>L3</u>
ALL	11 and glycosyltransferase	5	<u>L2</u>
ALL	sandrin.in.	94	<u>L1</u>